

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/601, 777 B
Source: IFW16
Date Processed by STIC: 12/14/2005

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 12/14/2005

PATENT APPLICATION: US/10/601,777B

TIME: 11:54:30

Input Set : A:\PTO.TS.txt

Output Set: N:\CRF4\12142005\J601777B.raw

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2 <110> APPLICANT: Okamoto, Tadashi
3     Takase, Hiromitsu
4     Hashimoto, Hiroyuki
W--> 5 <120> TITLE OF INVENTION: A METHOD FOR ACQUIRING INFORMATION OF
W--> 6     A BIOCHIP USING TIME OF FLIGHT SECONDARY ION
W--> 7     MASS SPECTROMETRY AND AN APPARATUS FOR ACQUIRING
W--> 8     INFORMATION FOR THE APPLICATION THEREOF
W--> 9 <130> FILE REFERENCE: 03560.003310
W--> 10 <140> CURRENT APPLICATION NUMBER: US/10/601,777B
11 <141> CURRENT FILING DATE: 2003-06-24
12 <150> PRIOR APPLICATION NUMBER: JP 2002-191208
13 <151> PRIOR FILING DATE: 2002-06-28
14 <150> PRIOR APPLICATION NUMBER: JP 2003-161862
15 <151> PRIOR FILING DATE: 2003-06-06
W--> 16 <160> NUMBER OF SEQ ID: 5
W--> 17 <210> SEQ ID NO: 1
18 <211> LENGTH: 40
19 <212> TYPE: DNA
20 <213> ORGANISM: Artificial sequence
W--> 21 <220> FEATURE:
22 <223> OTHER INFORMATION: Sample oligonucleotide
W--> 23 <400> SEQUENCE: 1
24 tttttttttt tttttttttt tttttttttt tttttttttt 40
25 <210> SEQ ID NO: 2
26 <211> LENGTH: 50
27 <212> TYPE: DNA
28 <213> ORGANISM: Artificial sequence
W--> 29 <220> FEATURE:
30 <223> OTHER INFORMATION: Sample oligonucleotide
W--> 31 <400> SEQUENCE: 2
32 tgcaggcatg caagcttggc actggccgtc gttttacaac gtcgtgactg 50
33 <210> SEQ ID NO: 3
34 <211> LENGTH: 20
35 <212> TYPE: RNA
36 <213> ORGANISM: Artificial sequence
W--> 37 <220> FEATURE:
38 <223> OTHER INFORMATION: Sample oligonucleotide
W--> 39 <400> SEQUENCE: 3
40 uuuuuuuuuu uuuuuuuuuu 20
41 <210> SEQ ID NO: 4
42 <211> LENGTH: 18
43 <212> TYPE: DNA
44 <213> ORGANISM: Artificial sequence

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W--> 45 <220> FEATURE:
46 <223> OTHER INFORMATION: Sample oligonucleotide
W--> 47 <400> SEQUENCE: 4
48 actggccgtc gttttaca 18
49 <210> SEQ ID NO: 5
50 <211> LENGTH: 50
51 <212> TYPE: DNA
52 <213> ORGANISM: Artificial sequence
W--> 53 <220> FEATURE:
54 <221> NAME/KEY: misc_feature
55 <222> LOCATION: (1)..(1)
56 <223> OTHER INFORMATION: Sample oligonucleotide
W--> 57 <220> FEATURE:
58 <221> NAME/KEY: misc_feature
59 <222> LOCATION: (1)..(1)
60 <223> OTHER INFORMATION: n is NCys-NH-(CH₂)₂-O-(CH₂)₂-O-CH₂CONH-T
W--> 61 <400> SEQUENCE: 5
W--> 62 ngcaggcatg caagcttggc actggccgtc gttttacaac gtcgtgactg 50

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/601,777B

DATE: 12/14/2005
TIME: 11:54:31

Input Set : A:\PTO.TS.txt
Output Set: N:\CRF4\12142005\J601777B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 1

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/601,777B

DATE: 12/14/2005

TIME: 11:54:31

Input Set : A:\PTO.TS.txt

Output Set: N:\CRF4\12142005\J601777B.raw

L:5 M:283 W: Missing Blank Line separator, <120> field identifier
L:9 M:283 W: Missing Blank Line separator, <130> field identifier
L:10 M:283 W: Missing Blank Line separator, <140> field identifier
L:16 M:283 W: Missing Blank Line separator, <160> field identifier
L:17 M:283 W: Missing Blank Line separator, <210> field identifier
L:21 M:283 W: Missing Blank Line separator, <220> field identifier
L:23 M:283 W: Missing Blank Line separator, <400> field identifier
L:29 M:283 W: Missing Blank Line separator, <220> field identifier
L:31 M:283 W: Missing Blank Line separator, <400> field identifier
L:37 M:283 W: Missing Blank Line separator, <220> field identifier
L:39 M:283 W: Missing Blank Line separator, <400> field identifier
L:45 M:283 W: Missing Blank Line separator, <220> field identifier
L:47 M:283 W: Missing Blank Line separator, <400> field identifier
L:53 M:283 W: Missing Blank Line separator, <220> field identifier
L:57 M:283 W: Missing Blank Line separator, <220> field identifier
L:61 M:283 W: Missing Blank Line separator, <400> field identifier
L:62 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0